SEQUENCE LISTING

<110> Remer, Ricardo A
Margis, Rogerio
Alves Ferreira, Marcio
Coronha Lima, Marcia

<120> Pharmaceutical Product Comprising Tissue of the Male Vegetal...

<130> 149456.00100

<140> 10/595610
<141> 2006-05-01

<150> PCT/BR04/00100
<151> 2004-11-12

<150> BR PI0305197-8
<151> 2003-11-13

<160> 3

<170> PatentIn version 3.3

<210> 1
<211> 1818
<212> DNA
<213> Artificial

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+ (5545-6757) - cDNA

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agaaataggt ttccttttct gtccctgtca cagagagagg gcagattttt tccttctcta 120

tctctttcag agagagatgg aagaaagttt tcttttctca gtatgttctc ttttctcatg

180

ccactgttgg aggttattaa gattattatt gcttctgtgg cctccgtaat cttcgtcggt
240

ttcgcctgtg taaccctcgc tggttctgcc gcagcattag tcgtaagcac cccggttttc
300

atcatattta gtcctgttct cgtaccagct acgatagcca cggttgtctt ggcgacagga
360

ttcacggccg gtggctcttt tggagcgacg gcacttggtc tcatcatgtg gcttgttaag
420

taagattatt ataacagctt atattgagat cactcgagat ttatgcttaa ttatataata
480

ttcataaacc tatagtttaa aagtatattg aacttcattt gttaacgtac tttataaata
540

ttgaacttcg ttcgttttct taattggtct ctaagtatat atacatactt ttttgtgtga
600

tgcagacgta ggatgggagt aaagccgaag gataatccac ctccggcagg acttccaccg
660

aattcgggag caggagcagg aggagctcaa agtctgatca aaaagtcaaa ggcaaagtct
720

aaaggtgggc ttaaggcttg gtgtaagaag atgttaaaaa gtaaattcgg tggtaaaaaa
780

ggcaagtccg ggggtggaaa aagtaaattt ggaggtaaag gcggtaagtc cgaaggtgaa
840

gaaggtatgt cgtctgggga tgaaggtatg tctggaagtg aaggaggtat gtccggaggt
900

gaaggaggta aatccaaaag tggaaaaggt aaactcaaag ctaaactcga aaagaaaaaa
960

ggtatgtccg gagggtccga gagtgaagaa ggtatgtctg gaagtgaagg aggtatgtct
1020

ggtggtggag gaagtaaatc caaaagtaaa aaaagtaaac tcaaagctaa attgggaaag
1080

aaaaaaggta tgtccggagg catgtcagga agtgaagaag gtatgtctgg aagtgaagga

1140

ggtatgtcca gtggtggagg aagtaaatcc aaaagtaaaa aaagtaaact caaagctaaa
1200

ttgggaaaga aaaaaggtat gtccggaggc atgtcaggaa gtgaagaagg tatgtctgga
1260

agtgaaggag gtatgtccgg aggtggagga ggtaaatcca aaagtagaaa aagtaaactc
1320

aaagctaaat tgggaaagaa aaaatgtatg tccggaggca tgtcaggaag tgaaggaggt
1380

atgtctggaa gtgaaggagg tatatccgga ggtggtatgt ctgggggcag tggaagtaaa
1440

cacaaaattg gaggaggtaa acacggaggt cttggaggta aattcggaaa gaaaagaggc
1500

atgtccggaa gtggaggagg catgtcagga agtgaaggag gtgtgtctgg aagtgaagga
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1620

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1680

ggtatgtctg gaagtgaagg aggtatgtcg gaaagtggta tgtccgggag tggaggggggt
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Glu Gly Arg Phe Phe Pro Ser Leu Ser Leu Ser Glu Arg Asp Gly Arg
        35                  40                  45

Lys Phe Ser Phe Leu Ser Met Phe Ser Phe Leu Met Pro Leu Leu Glu
    50                  55                  60

Val Ile Lys Ile Ile Ile Ala Ser Val Ala Ser Val Ile Phe Val Gly
65                  70                  75                  80

Phe Ala Cys Val Thr Leu Ala Gly Ser Ala Ala Ala Leu Val Val Ser
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Thr Pro Val Phe Ile Ile Phe Ser Pro Val Leu Val Pro Ala Thr Ile
            100                 105                 110

Ala Thr Val Val Leu Ala Thr Gly Phe Thr Ala Gly Gly Ser Phe Gly
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Ala Thr Ala Leu Gly Leu Ile Met Trp Leu Val Lys Arg Arg Met Gly
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Val Lys Pro Lys Asp Asn Pro Pro Pro Ala Gly Leu Pro Pro Asn Ser
145                 150                 155                 160

Gly Ala Gly Ala Gly Gly Ala Gln Ser Leu Ile Lys Lys Ser Lys Ala
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Lys Ser Lys Gly Gly Leu Lys Ala Trp Cys Lys Lys Met Leu Lys Ser
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Lys Phe Gly Gly Lys Lys Gly Lys Ser Gly Gly Gly Lys Ser Lys Phe
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Gly Gly Lys Gly Gly Lys Ser Glu Gly Glu Glu Gly Met Ser Ser Gly
210 215 220

Asp Glu Gly Met Ser Gly Ser Glu Gly Gly Met Ser Gly Gly Glu Gly
225 230 235 240

Gly Lys Ser Lys Ser Gly Lys Gly Lys Leu Lys Ala Lys Leu Glu Lys
245 250 255

Lys Lys Gly Met Ser Gly Gly Ser Glu Ser Glu Glu Gly Met Ser Gly
260 265 270

Ser Glu Gly Gly Met Ser Gly Gly Gly Gly Ser Lys Ser Lys Ser Lys
275 280 285

Lys Ser Lys Leu Lys Ala Lys Leu Gly Lys Lys Lys Gly Met Ser Gly
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Gly Met Ser Gly Ser Glu Glu Gly Met Ser Gly Ser Glu Gly Gly Met
305 310 315 320

Ser Ser Gly Gly Gly Ser Lys Ser Lys Ser Lys Lys Ser Lys Leu Lys
325 330 335

Ala Lys Leu Gly Lys Lys Lys Gly Met Ser Gly Gly Met Ser Gly Ser
340 345 350

Glu Glu Gly Met Ser Gly Ser Glu Gly Gly Met Ser Gly Gly Gly Gly
355 360 365

Gly Lys Ser Lys Ser Arg Lys Ser Lys Leu Lys Ala Lys Leu Gly Lys
370 375 380

Lys Lys Cys Met Ser Gly Gly Met Ser Gly Ser Glu Gly Gly Met Ser
385 390 395 400

Gly Ser Glu Gly Gly Ile Ser Gly Gly Gly Met Ser Gly Gly Ser Gly
405 410 415

Ser Lys His Lys Ile Gly Gly Gly Lys His Gly Gly Leu Gly Gly Lys
420 425 430

Phe Gly Lys Lys Arg Gly Met Ser Gly Ser Gly Gly Gly Met Ser Gly
435 440 445

Ser Glu Gly Gly Val Ser Gly Ser Glu Gly Ser Met Ser Gly Gly Gly
450 455 460

Met Ser Gly Gly Ser Gly Ser Lys His Lys Ile Gly Gly Gly Lys His
465 470 475 480

Gly Gly Leu Arg Gly Lys Phe Gly Lys Lys Arg Gly Met Ser Gly Ser
485 490 495

Glu Gly Gly Met Ser Gly Ser Glu Gly Gly Met Ser Glu Ser Gly Met
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<223> Complete sequence of the promoter region of the AtGRP17

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120

attcaatcgt gagacattga aattgtcgtt tctccattac ctttttggaa gaaaaaccat
180

cgaaagctag ctaagacttt ttttattaaa cgaacttgct actatttcta tgttttcttt
240

gaaatgaaaa ttaaatttgt tactgtttca cctaaaactc aaaagtattg ctttttaatt
300

ttattattaa gaaaaactaa tcttatttat gttaagaaac ctgtcaattt ttcattgtta
360

atttcggctc tataattatt aattaacaat caatttctca aaaattgcaa tcatgattat
420

gattagatat atattagttg gattgtgatg catttttgt aatataaaat ggatgtttgt
480

attagtttct cactcatgta attaaacacc aaatgctaga aactagtact tttgtttctc
540

agctctcgtc tattgttata tctgcaacac gaacaaaaac cttatctagg tgttatatat
600

cacggttatg tttatgagtt agaagggatt cttcaacaaa aatcacggaa ctacttgtat
660

atatgtatgt gtgtatccga tcgaggttga cttccggggt tggacgttga agaagacgaa
720

ttcattgatt gggcttatat atgggcatgt attacttggt tcaagtttgt aacactttta
780

gctttttcaa ttctattcga aaccaaaata ttgggctata tatctttata caaccttcaa
840

gataaattgg accaatttta gaagagcaaa ttgaacccgg ccgttagcgt tagccaaacc
900

ccaactcctt ttcagtacaa ttaaatcaag aatttctaat aaatcgtgaa tttctagaca

960

tacatatcat aatttcgtca aagcgagcct acacctagtt ttgagctaca taactctttt
1020

ctttttttt ttatgattag gaggtttcaa aacccttgga cccataattt cttataatta
1080

gttttgtaat actaaattta ccattgagag cgacctctcg tcactagtaa ttcgaagatc
1140

tcatattcat gacctatatt aaccatcttc cagtcaagta atttcaatcg aaattcatca
1200

aaatcatata tttaacttag taatcacata tgatatggct aatatacgta atataacgat
1260

aaagatttct tcacgctttg atattccata aagcaatgga aatatggaat ggaagaaaac
1320

atttgaattt tacaagaaac aataaataga aggcctacaa aacatgacaa cccacacaca
1380

cacacacgaa aagagaaaat ataaagaagg acatgtaacg tgacgtagcg tagatctcca
1440

ttcactccaa tcgttttgca tggagcatgc atgtgtgtgt accgtgcacg tagtagagac
1500

cacacaactc cttcataaaa gccctctctc tcttaccatc accaaaacac aacaatccga
1560

tcagaaaat
1569